

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/586, 403
Source: IFWP
Date Processed by STIC: 07/26/2006

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 07/26/2006

PATENT APPLICATION: US/10/586,403

TIME: 14:18:54

Input Set : A:\00786.455003.SEQLIST.TXT

Output Set: N:\CRF4\07262006\J586403.raw

4 <110> APPLICANT: RAHME, Laurence
5 DEZIEL, Eric
6 LEPIINE, Francois
7 TOMPKINS, Ronald G.
8 XIAO, Gaoping
10 <120> TITLE OF INVENTION: Methods For Identifying Candidate
11 Compounds For Treating, Reducing, or Preventing Pathogenic
12 Infections
14 <130> FILE REFERENCE: 00786/455003
C--> 16 <140> CURRENT APPLICATION NUMBER: US/10/586,403
C--> 16 <141> CURRENT FILING DATE: 2006-07-18
16 <150> PRIOR APPLICATION NUMBER: PCT/US05/02174
17 <151> PRIOR FILING DATE: 2005-01-21
19 <150> PRIOR APPLICATION NUMBER: US 60/538,361
20 <151> PRIOR FILING DATE: 2004-01-22
22 <150> PRIOR APPLICATION NUMBER: US 60/538,278
23 <151> PRIOR FILING DATE: 2004-01-21
25 <160> NUMBER OF SEQ ID NOS: 17
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30 <211> LENGTH: 1554
31 <212> TYPE: DNA
32 <213> ORGANISM: Pseudomonas aeruginosa
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36 gccgtttatc actatcgggg ccagactctc agccggctgc aatgcccgc ac ctacattctc 120
37 tcccaggcca gccaactggc cccgctgttc aagccggcg atcgcgttgt gctggcggttg 180
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40 gccagcctgg tggtgcgtga agccgatgca ccgtcgctga gcggtccctt ggccgcgttg 360
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46 tggttcagcg gaggcctcgcc gctgctcgac gataacctggc cgagccccga gcgggttctg 720
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49 ccgctgcgcg cggcgagtt cgaattctgg gccgcgcacg ggctggagat ctgcgcacggc 900
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52 actatcgagg aagcggggccg gcaaggcgtg ctgttgggtgc gtggcccagg gctgagtcgg 1080
53 ggttaactggc gggccagcga agagcagcag ggcgcgttc caggtggctg gtaccgcacc 1140

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 55 ctgttcaagg tgaatggccg ctgggtggtg ccgaccagg tcgagcagc gatctgccgt 1260
 56 catctgccgg aagtgagcga ggcggtctg gttcctacct gccggctgca cgacggcttg 1320
 57 cgtccgaccc tggtcgtcac cctggccact ccgctggacg acaaccagat cctgctggcg 1380
 58 cagcgcatcg accagcatct cgccgaacag attccctcgc acatgctgcc cagccaattg 1440
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 87 <213> ORGANISM: Pseudomonas aeruginosa
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 92 gtcaatcgcc ggggttattt cgacccgcgg aacggcgaga acgagttcag cctggtggtc 180
 93 cggggccggc agcgccgtct gctgtacgc gataccgcgc cggatagcgt ggacatgtc 240
 94 atctgttccgg ctccctcgcc gatcatgacc gacgcggcg atgtcttgcg ggacactgcgc 300
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 101 gccaaggccgc ggctttattt ctgcgttgcg acgacggcc agaacaagat ggccagcttc 720
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 103 gatgacatcg attatttcgt ctccaccag ccagcgccgt tcctggtcaa ggcttggcc 840
 104 gagggcatcg gtgcccgtcc tgagcgtac caactgacga tgggcgatc cggcgtatg 900
 105 atctccgttt ccatcccgtt caccctgtg accggccgtc gcgaggccaa gatccggcccc 960
 106 ggcgatcgta tcgtcatggc cggcgccagcc actggctggg ggttcgcgcg ccaggtctgg 1020

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116 catgacacctgg tagggcgcat caatacgtcg gacgagttca tcgtcgaacg taccggcgtg 120
117 cgcacccgcgt atcacgtcg a cccggaaacag gcggtcagcg cgctgatggt gccggcggcg 180
118 cgccaggccca tcgaggtcg cgggctgtg ccggaggaca tcgacctgtt gctggtaac 240
119 acctgtcgcc cgaccacca cgaccgtcc caggcctgcc tgatccagcc gctgctgggc 300
120 ctgcggcaca tcccggtaact ggatatccgg gcacagtca gcgggttgcgt gtacggcttg 360
121 cagatggctc gccccagat cctcgccggg ctggcacggc atgtcctggt ggtctgcggc 420
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124 gacctgcgc tggggccgca cggcaactac ttgcacctgc tgatgaccgc ggcggccgggt 600
125 agtgcctcgcc cgaccccttccct cgacgagaat gtcctgcgcg a gggcgccccg cgagttccctc 660
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129 gtgaccgtgg atcgctctggg caacatggct tcggcctcgca ccccggtcac gctggcgatg 900
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143 ctgtgtccgc ggctggccaa cgtacaggc tcggcgtccg agcggacctg ccaggccctgg 300
144 aagtccggaaa ggcggtgcg ggtggtcagc cgcttgaacc ggcaactgtt gcgtgcggag 360
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152 ctgagcgagg agctgcaccc cccctggggt gggcagagcg tcgacttccct gcccggcgaa 840
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154 gactga          906
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158 <212> TYPE: DNA
159 <213> ORGANISM: Pseudomonas aeruginosa
161 <400> SEQUENCE: 6

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 219 20 25 30
 220 Leu Gln Cys Arg Thr Tyr Ile Leu Ser Gln Ala Ser Gln Leu Ala Arg
 221 35 40 45
 222 Leu Leu Lys Pro Gly Asp Arg Val Val Leu Ala Leu Asn Asp Ser Pro
 223 50 55 60
 224 Ser Leu Ala Cys Leu Phe Leu Ala Cys Ile Ala Val Gly Ala Ile Pro
 225 65 70 75 80
 226 Ala Val Ile Asn Pro Lys Ser Arg Glu Gln Ala Leu Ala Asp Ile Ala
 227 85 90 95
 228 Ala Asp Cys Gln Ala Ser Leu Val Val Arg Glu Ala Asp Ala Pro Ser
 229 100 105 110
 230 Leu Ser Gly Pro Leu Ala Pro Leu Thr Leu Arg Ala Ala Gly Arg
 231 115 120 125
 232 Pro Leu Leu Asp Asp Phe Ser Leu Asp Ala Leu Val Gly Pro Ala Asp
 233 130 135 140
 234 Leu Asp Trp Ser Ala Phe His Arg Gln Asp Pro Ala Ala Ala Cys Phe
 235 145 150 155 160
 236 Leu Gln Tyr Thr Ser Gly Ser Thr Gly Ala Pro Lys Gly Val Met His
 237 165 170 175
 238 Ser Leu Arg Asn Thr Leu Gly Phe Cys Arg Ala Phe Ala Thr Glu Leu
 239 180 185 190
 240 Leu Ala Leu Gln Ala Gly Asp Arg Leu Tyr Ser Ile Pro Lys Met Phe
 241 195 200 205
 242 Phe Gly Tyr Gly Met Gly Asn Ser Leu Phe Phe Pro Trp Phe Ser Gly
 243 210 215 220
 244 Ala Ser Ala Leu Leu Asp Asp Thr Trp Pro Ser Pro Glu Arg Val Leu
 245 225 230 235 240
 246 Glu Asn Leu Val Ala Phe Arg Pro Arg Val Leu Phe Gly Val Pro Ala
 247 245 250 255
 248 Ile Tyr Ala Ser Leu Arg Pro Gln Ala Arg Glu Leu Leu Ser Ser Val
 249 260 265 270
 250 Arg Leu Ala Phe Ser Ala Gly Ser Pro Leu Pro Arg Gly Glu Phe Glu
 251 275 280 285
 252 Phe Trp Ala Ala His Gly Leu Glu Ile Cys Asp Gly Ile Gly Ala Thr
 253 290 295 300
 254 Glu Val Gly His Val Phe Leu Ala Asn Arg Pro Gly Gln Ala Arg Ala
 255 305 310 315 320
 256 Asp Ser Thr Gly Leu Pro Leu Pro Gly Tyr Glu Cys Arg Leu Val Asp
 257 325 330 335
 258 Arg Glu Gly His Thr Ile Glu Glu Ala Gly Arg Gln Gly Val Leu Leu
 259 340 345 350
 260 Val Arg Gly Pro Gly Leu Ser Pro Gly Tyr Trp Arg Ala Ser Glu Glu
 261 355 360 365
 262 Gln Gln Ala Arg Phe Ala Gly Gly Trp Tyr Arg Thr Gly Asp Leu Phe
 263 370 375 380

VERIFICATION SUMMARY

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L:16 M:270 C: Current Application Number differs, Replaced Current Application No
L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date